l N	umber: 0918-6/6/2	CRF Process Edited by:	Ing Date: 1	
	Changed a file from non-ASCII to ASCII	Verified by: _		(STIC s
	Changed the margins in cases where the sequence text was "wrapped to the County Application Data section, specifically	-MT"		in T
	Edited a format error in the Current Application Data section, specifical	ly:		
	Edited the Current Application Data section with the actual current num applicant was the prior application data; or other	ber. The nur	mber inputt	ed by the
	Added the mandatory heading and subheadings for *Current Application	n Data*.		
	Edited the "Number of Sequences" field. The applicant spelled out a nu	umber instead	d of using a	ກ integer.
(Changed the spelling of a mandatory field (the headings or subheading	s), specificall	У	
(Corrected the SEQ ID NO when obviously incorrect. The sequence nur	mbers that we	ere edited v	vere:
ı	Inserted or corrected a nucleic number at the end of a nucleic line. SEC	Q ID NO's ed	lited:	
(a	Corrected subheading placement. All responses must be on the same I applicant placed a response below the subheading, this was moved to it	ine as each s Is appropriate	subheading o place.	. If the
	Inserted colons after headings/subheadings. Headings edited included	:		
l	Deleted extra, inválid, headings used by an applicant, specifically:			
	Deleted: non-ASCII "garbage" at the beginning/end of files; se page numbers throughout text; other invalid text, such as	cretary initial	s/filename	at end of file
	Inserted mandatory headings, specifically:		<u> </u>	
	Corrected an obvious error in the response, specifically:			
	Edited identifiers where upper case is used but lower case is required, or	or vice versa.		
(Corrected an error in the Number of Sequences field, specifically:			
_	A "Hard Page Break" code was inserted by the applicant. All occurrence	es had to be	deleted.	
	Peleted ending stop codon in amino acid sequences and adjusted the *ue to a Patentin bug). Sequences corrected:			ingly (error
_	Other: For Sign 90491 mount agail 409012, respectively to Due of complete, mounterpretation	in 22237	7, to prev	, y line

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

Peter

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PCT09

RAW SEQUENCE LISTING DATE: 11/07/2001 PATENT APPLICATION: US/09/856,662 TIME: 09:18:51

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

- 3 <110> APPLICANT: MORIBE, Toyoki et al.
 5 <120> TITLE OF INVENTION: Method for typing HLA class 1 genes
 7 <130> FILE REFERENCE: 0032-0261P
 9 <140> CURRENT APPLICATION NUMBER: US 09/856,662
 10 <141> CURRENT FILING DATE: 2001-05-24
 12 <150> PRIOR APPLICATION NUMBER: JP P1998-335151
 13 <151> PRIOR FILING DATE: 1998-11-26
 15 <160> NUMBER OF SEQ ID NOS: 130
 17 <170> SOFTWARE: Patentin Ver. 2.0
- 19 <210> SEQ ID NO: 1 20 <211> LENGTH: 23
- 21 <212> TYPE: DNA
- 22 <213> ORGANISM: Artificial Sequence
- 24 <220> FEATURE:
- 25 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA probe A98T
- 27 <400> SEQUENCE: 1
- 28 gaggtatttc ttcacatccg tgt
- 30 <210> SEQ ID NO: 2
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- 32 <212> TYPE: DNA
- 33 <213> ORGANISM: Artificial Sequence
- 35 <220> FEATURE:
- 36 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA probe A98A
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- 39 atgaggtatt tctacacctc cgtgt
- 41 <210> SEQ ID NO: 3
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- 43 <212> TYPE: DNA
- 44 <213> ORGANISM: Artificial Sequence
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- 50 tacgtggaca acacgcagt
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- 54 <212> TYPE: DNA
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- 65 <212> TYPE: DNA
- 66 <213> ORGANISM: Artificial Sequence
- 68 <220> FEATURE:
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RAW SEQUENCE LISTING

DATE: 11/07/2001

PATENT APPLICATION: US/09/856,662

TIME: 09:18:51

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- 71 <400> SEQUENCE: 5
- 72 caggagaggc ctgag

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- 88 <213> ORGANISM: Artificial Sequence
- 90 <220> FEATURE:
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- 94 <400> SEQUENCE: 7
- 95 ttgggacctg cagacacg
- 97 <210> SEQ ID NO: 8
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- 102 <220> FEATURE:
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 - 109 <210> SEQ ID NO: 9
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- 114 <220> FEATURE:
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- 117 <400> SEQUENCE: 9
- 118 gacacggaat gtgaaggc 120 <210> SEQ ID NO: 10
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- 122 <212> TYPE: DNA
- 123 <213> ORGANISM: Artificial Sequence
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- 129 tgaaggccca ctcacagact
- 131 <210> SEQ ID NO: 11
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- 134 <213> ORGANISM: Artificial Sequence
- 136 <220> FEATURE:

RAW SEQUENCE LISTING DATE: 11/07/2001 PATENT APPLICATION: US/09/856,662 TIME: 09:18:51

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

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202 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING

DATE: 11/07/2001

PATENT APPLICATION: US/09/856,662

TIME: 09:18:51

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

- 204 <220> FEATURE:
- 205 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA probe A368A
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- 208 gaggatgtat ggctgc

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- 210 <210> SEQ ID NO: 18
- 211 <211> LENGTH: 16
- 212 <212> TYPE: DNA
- 213 <213> ORGANISM: Artificial Sequence
- 215 <220> FEATURE:
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- 218 <400> SEQUENCE: 18
- 219 gaggatgtgt ggctgc
- 221 <210> SEQ ID NO: 19
- 222 <211> LENGTH: 16
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- 229 <400> SEQUENCE: 19
- 230 gaggatgttt ggctgc
- 16 232 <210> SEQ ID NO: 20
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- 234 <212> TYPE: DNA
- 235 <213> ORGANISM: Artificial Sequence
- 237 <220> FEATURE:
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- 241 cgcttcctgc gcgggt
- 243 <210> SEQ ID NO: 21
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- 245 <212> TYPE: DNA
- 246 <213> ORGANISM: Artificial Sequence
- 248 <220> FEATURE:
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- 251 <400> SEQUENCE: 21
- 252 caggacgett acgacgg
- 254 <210> SEQ ID NO: 22
- 255 <211> LENGTH: 16
- 256 <212> TYPE: DNA
- 257 <213> ORGANISM: Artificial Sequence
- 259 <220> FEATURE:
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- 262 <400> SEQUENCE: 22
- 265 <210> SEQ ID NO: 23
- 16 263 catcgccctg aacgag
- 266 <211> LENGTH: 16
- 267 <212> TYPE: DNA
- 268 <213> ORGANISM: Artificial Sequence
- 270 <220> FEATURE:

RAW SEQUENCE LISTING

DATE: 11/07/2001

PATENT APPLICATION: US/09/856,662

TIME: 09:18:51

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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\I856662.raw

- 271 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA probe A485A
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- 274 gcggacaagg cagete
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- 276 <210> SEQ ID NO: 24
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- 279 <213> ORGANISM: Artificial Sequence
- 281 <220> FEATURE:
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- 284 <400> SEQUENCE: 24
- 285 geggeeegtg tggegg
- 287 <210> SEQ ID NO: 25
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- 289 <212> TYPE: DNA
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- 296 cggccgttg ggcggag
- 298 <210> SEO ID NO: 26
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- 319 gagcagcgga gagtc
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- 330 gagcagcaga gagcct 332 <210> SEQ ID NO: 29
- 333 <211> LENGTH: 15
- 334 <212> TYPE: DNA
- 335 <213> ORGANISM: Artificial Sequence
- 337 <220> FEATURE:

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/856,662 TIME: 09:18:52

DATE: 11/07/2001

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\11072001\1856662.raw

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RAW SEQUENCE LISTING
                                                              DATE: 10/29/2001
                     PATENT APPLICATION: US/09/856,662
                                                              TIME: 13:07:49
                                                                     Done H
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                     Output Set: N:\CRF3\10292001\I856662.raw
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      3 <110 > APPLICANT: MORIBE, Toyoki et al.
      5 <120> TITLE OF INVENTION: Method for typing HLA class 1 genes
      7 -: 130> FILE REFERENCE: 0032-0261P
      9 <140 > CURRENT APPLICATION NUMBER: US 09/856,662
     10 <141> CURRENT FILING DATE: 2001-05-24
     12 <150> PRIOR APPLICATION NUMBER: JP P1998-335151
     13 <151> PRIOR FILING DATE: 1998-11-26
                                                                                        - 1.40 J
     15 <160> NUMBER OF SEO ID NOS: 130
     17 <170> SOFTWARE: PatentIn Ver. 2.0
                                                              Cestion compa
ERRORED SEQUENCES
                                                              JAMESTO CHERRETTE NEEDLES
     1023 <210> SEQ ID NO: 90
     1024 <211> LENGTH: 19
     1025 +212> TYPE: DNA
     1026 <213> ORGANISM: Artificial Sequence
     1028 <220> FEATURE:
     1029 <223> OTHER INFORMATION: Description of Artificial Sequence: PCR primer \mathcal{E}_{T}
              (cga011)
E--> 1030
                               invocal could be on to fine sabore to sound computer my nature producing
W--> 1032 <210> SEQ ID NO:
W--> 1032 <211> LENGTH:
W--> 1032 <212> TYPE:
W--> 1032 <213> ORGANISM:
     1032 -: 400> SEQUENCE: 90
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     1037 +212> TYPE: DNA
     1038 <213> ORGANISM: Artificial Sequence
     1040 <2220> FEATURE:
     1041 +223 OTHER INFORMATION: Description of Artificial Sequence: PCR primer \epsilon
E--> 1042
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                                        - Same Bo Blocke
W--> 1044 <210> SEQ ID NO:
W--> 1044 <211> LENGTH:
W--> 1044 <212> TYPE:
W--> 1044 <213> ORGANISM:
    1044 <400> SEQUENCE: 91
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/856,662

DATE: 10/29/2001 TIME: 13:07:51

Input Set : A:\0032-0261P.ST25.txt

Output Set: N:\CRF3\10292001\1856662.raw

L:1030 M:254 E: No. of Bases conflict, LENGTH:Input:11 Counted:3 SEQ:90 L:1030 M:112 C: (48) String data converted to lower case,
L:1030 M:252 E: No. of Seq. differs, -211>LENGTH:Input:19 Found:3 SEQ:90 L:1032 M:282 W: Numeric Field Identifier Missing, <210> is required.
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L:1042 M:254 E: No. of Bases conflict, LENGTH:Input:12 Counted:3 SEQ:91 M:112 Repeated in SeqNo=91
L:1042 M:252 E: No. of Seq. differs, <211>LENGTH:Input:19 Found:3 SEQ:91 L:1044 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:1044 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:1044 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:1044 M:282 W: Numeric Field Identifier Missing, <212> is required.